

Figure 1

Sequence Name:

GW.S.ctg16335-000003.31.0

Figure 1A

LLAPTGSLFRNCTQDGWSETFPRPNLACGVNVNDSSNEKRSYLLKLKVMYTVGYSSSLVM
LLVALGILCAFRRLHCTRNYIHMHLFVSFILRALSNFIKDAVLFSSDDVTYCDAHRGCKL
VMVLFXYCIMANYSWLLVEGSTFTHxLAISFFSERKYLQGFVAFGWGSPAIFVALWAIAR
HFLEDVGCWDINANASIWWIIRGPVILSILNFILFINILRILMRKLRTQETRGNEVSHYK
RLARSTLLLIPLFGIHYIVFAFSPEDAMEIQLFF

Figure 1B

CTCTTGGCACCCACAGGTTCTTGTTCGGAACTGCACACAGGATGGCTGGTCAGAAACC
TTCCCCAGGCCTAATCTGGCCTGTGGCGTTAATGTGAACGACTCTTCCAACGAGAAGCGG
CACTCCTACCTGCTGAAGCTGAAAGTCATGTACACCGTGGGCTACAGCTCCTCCCTGGTC
ATGCTCCTGGTCGCCCTTGGCATCCTCTGTGCTTTCCGGAGGCTCCACTGCACTCGCAAC
TACATCCACATGCACCTGTTTCGTGTCCTTCATCCTTCGTGCCCTGTCCAACCTCATCAAG
GACGCCGTGCTCTTCTCCTCAGATGATGTACCTACTGCGATGCCACAGGGCGGGCTGC
AAGCTGGTCATGGTGCTGTTC

Figure 1C

TACTGCATCATGGCCAACTACTCCTGGCTGCTGGTGGAAGGCTCTACCTTCACACATNTC
CTCGCCATCTCCTTCTTCTCTGAAAGAAAGTACCTCCAGGGATTTGTGGCATTTCGGATGG
GGTTCTCCAGCCATTTTTGTGTGCTTTGTGGGCTATTGCCAGACACTTTCTGGAAGATGTT
GGGTGCTGGGACATCAATGCCAACGCATCCATCTGGTGATCATTTCGTGGTCCTGTGATC
CTCTCCATCCTGATTAATTTTCATCCTTTTCATAAACATTCTAAGAATCCTGATGAGAAAA
CTTAGAACCCAAAGAAACAAGAGGAAATGAAGTCAGCCATTATAAGCGCCTGGCCAGGTCC
ACTCTCCTGCTGATCCCCCTCTTTGGCATCCACTACATCGTCTTCGCCTTCTCCCCAGAG
GACGCTATGGAGATCCAGCTGTTTTTT

Figure 1D

CTCTTGGCACCCACAGGTTCTTGTTCGGAACTGCACACAGGATGGCTGGTCAGAAACC
TTCCCCAGGCCTAATCTGGCCTGTGGCGTTAATGTGAACGACTCTTCCAACGAGAAGCGG
CACTCCTACCTGCTGAAGCTGAAAGTCATGTACACCGTGGGCTACAGCTCCTCCCTGGTC
ATGCTCCTGGTCGCCCTTGGCATCCTCTGTGCTTTCCGGAGGCTCCACTGCACTCGCAAC
TACATCCACATGCACCTGTTTCGTGTCCTTCATCCTTCGTGCCCTGTCCAACCTCATCAAG
GACGCCGTGCTCTTCTCCTCAGATGATGTACCTACTGCGATGCCACAGGGCGGGCTGC
AAGCTGGTCATGGTGCTGTTC
TACTGCATCATGGCCAACTACTCCTGGCTGCTGGTGGAAGGCTCTACCTTCACACATNTC
CTCGCCATCTCCTTCTTCTCTGAAAGAAAGTACCTCCAGGGATTTGTGGCATTTCGGATGG
GGTTCTCCAGCCATTTTTGTGTGCTTTGTGGGCTATTGCCAGACACTTTCTGGAAGATGTT
GGGTGCTGGGACATCAATGCCAACGCATCCATCTGGTGATCATTTCGTGGTCCTGTGATC
CTCTCCATCCTGATTAATTTTCATCCTTTTCATAAACATTCTAAGAATCCTGATGAGAAAA
CTTAGAACCCAAAGAAACAAGAGGAAATGAAGTCAGCCATTATAAGCGCCTGGCCAGGTCC
ACTCTCCTGCTGATCCCCCTCTTTGGCATCCACTACATCGTCTTCGCCTTCTCCCCAGAG
GACGCTATGGAGATCCAGCTGTTTTTT

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Figure 2A

Figure 2B

CCCACTTTTCATACTATTCTCTTTCCAGCCTGGTGACAAGAGAACAAAACATATCTGTGTCTACTGGGAGGGATCAGAGGGAGGCCACTGGTCCACGGAGGGCTGCTCTCATGTGCACAGCAACGGTTCCTTACACCAAAATGCAAGTGCTTCCATCTGTCCAGCTTTGCCGTCTCTGTGGCTCTTGCCCCCAAGGAGGACCTGTGCTGACCGTGATACCCAGGTGGGGCTGACCATCTCTCTGCTGTGCCTCTTCTGGCCATCCTCACCTTCTCTGTGCCGGCCCATCCAGAACACCAGCACCTCCCTCCATCTAGAGCTCTCCCTCTGCCTCTTCTGGCCCCACCTCCTGTTCCCTGACGGGCATCAACAGAACTGAGCCTGAGGTGCTGTGCTCCATCATTGCAGGGCTGCTGCAC TTCTCTACCTGGCTTGCTTACCTGGATGCTCCTGGAAGGGCTGCACCTCTTCCCTCACC GTCAGGAACCTCAAGGTGGCCAACTACACCAGCACGGGCAGATTCAAGAAGAGGTTTCATGTACCTGTAGGCTACGGGATCCCAGCTGTGATTATTGCTGTGTGAGCAATAGTTGGACCC CAGAATTATGGAACATTTACTACTGTTGGCTCAAGCTTGATAAAGGATTCATCTGGAGC TTCATGGGGCCAGTAGCAGTCATTATCTTGATAAACCTGGTGTTCTACTTCCAAGTTCTGTGAGATTTTGAGAAGCAAACCTTCTCCCTCAATAAAGAAGTTTCCACCATTGAGGACACC AGAGTCATGACATTTAAAGCCATTTCTCAGCTATTTATCCTGGGCTGTTCTTGGGGCCCTT GGTTTTTTTATGTTGAAGAAGTAGGGAAGACGATTGGATCAATCATTGCATACTCATTG ACCATCATCAACACCCTTCAGGGAGTGTTGCTCTTTGTGGTACACTGCTCCTTAATCGC CAGGTAAGG

Figure 3

Sequence Name:

GW.S.ctgl3100-000000.33.0

Figure 3A

QHSDAVHDLDDLVDITWVGILLSLVCLLICIFTFCFFRGLQSDRNTIHKNLCSLFFVAELL
FLIGINRTDQPACAVFAALLHFFFFLAAFTWMFLEGVQLYIMLVEVFESEHSRRKYFYLVG
YGMPALIVAVSAAVDYRSYGTDKVCWLRLDTYFIWSFIGPATLIIMNVIFLGIALYKMFH
HTAILKPESGCLDNIKLKINIPIIKSIYIYMYICMCV

Figure 3B

CAGCACAGTGATGCGGTCCATGACCTCCTTCTGGATGTGATCACGTGGGTGGAATTTTG
CTGTCCCTTGTTTGTCTCCTGATTTGCATCTTCACATTTTGCTTTTTCCGGGGGCTCCAG
AGTGACCGTAACACCATCCACAAGAACCTCTGCATCAGTCTCTTTGTAGCAGAGCTGCTC
TTCCTGATTGGGATCAACCGAACTGACCAACCAATTGCCTGTGCTGTTTTCGCTGCCCTG
TTACATTTCTTCTTCTTGGCTGCCTTCACCTGGATGTTCTGGAGGGGGTGCAGCTTTAT
ATCATGCTGGTGGAGGTTTTTGAGAGTGAACATTCACGTAGGAAATACTTTTATCTGGTC
GGCTATGGGATGCCTGCACTCATTGTGGCTGTGTCAGCTGCAGTAGACTACAGGAGTTAT
GGAACAGATAAAGTATGTTGGCTCCGACTTGACACCTACTTCATTTGGAGTTTTATAGGA
CCAGCAACTTTGATAATTATGCTTAATGTAATCTTCCTTGGGATTGCTTTATATAAAATG
TTTCATCATACTGCTATACTGAAACCTGAATCAGGCTGTCTTGATAACATCAAGTTAAAA
ATTAATATTCCAATTATAAAATCTATTTATATCTATATGTATATATGCATGTGTGTG

Figure 3A

Figure 4

Sequence Name:

GW.A.ctg12444-000001.0.2

Figure 4A

GNVAVAFVYYKSIGPLLSSSDNFLKPNQYDNSEEEERVISSVISVSMSSNPPTLYELEK
ITFTLSHRKTDYRSLCAFWNYSPTMNGSWSSSEGCELTYSNETHTSCRCNHLTHFAILM
SSGPSIIKDYNILTRITQLGIIISLICLAICIFTFWFFSEIQSTRTTIHKNLCCSLFLAE
LVFLVGINTNTNKFCIIAGLLHYFFLAFAWMCIEGHLHYLIVVGVIYNKGFLHKNFYI
FGYLSPAVVVGFSALGYRYYGTTKVCWLSTENNFIWSFIGPACLIILVCIYKIVITIQK
SDDH

Figure 4B

GGCAATGTTGCAGTTGCATTTGTATATTATAAGAGTATTGGTCCTTTGCTTTCATCATCT
GACAACTTCTTATTGAAACCTCAAAATTATGATAATTCTGAAGAGGAGGAAAGAGTCATA
TCTTCAGTAATTTTCAGTCTCAATGAGCTCAAACCCACCCACATTATATGAAC TTGAAAA
ATAACATTTACATTAAAGTCATCGAAAGGTACAGATAGGTATAGGAGTCTATGTGCATTT
TGGAATTACTCACCTGATACCATGAATGGCAGCTGGTCTTCAGAGGGCTGTGAGCTGACA
TACTCAAATGAGACCCACACCTCATGCCGCTGTAATCACCTGACACATTTTGCAATTTTG
ATGTCCTCTGGTCCTTCCATTGGTATTAAAGATTATAATATTCTTACAAGGATCACTCAA
CTAGGAATAATTATTTCACTGATTTGTCTTGCCATATGCATTTTTACCTTCTGGTTCTTC
AGTGAAATTCAAAGCACCAGGACAACAATTCACAAAAATCTTTGCTGTAGCCTATTTCTT
GCTGAACTTGTTTTCTTGTTGGGATCAATAACAATACTAATAAGCTCTTCTGTTCAATC
ATTGCCGGACTGCTACACTACTTCTTTTTAGCTGCTTTTGCATGGATGTGCATTGAAGGC
ATACATCTCTATCTCATTGTTGTGGGTGTCATCTACAACAAGGGATTTTTGCACAAGAAT
TTTTATATCTTTGGCTATCTAAGCCCAGCCGTGGTAGTTGGATTTTCGGCAGCACTAGGA
TACAGATATTATGGCACAACCAAAGTATGTTGGCTTAGCACCGAAAACAAC TTTATTGG
AGTTTTATAGGACCAGCATGCCTAATCATTCTTGTATGTATATATAAAATTGTTATTACA
ATTCAAAAAAGTGATGATCAT

Figure 4B "Sequence"

Figure 5

Sequence Name:

GW.S.ctg12789-000004.100.0

Figure 5A

GAWATTGCSVAALYLDSTACFCNHSTSFAILLQIYEVQGP EEESLLRTL SFVGC GVSFCA
LTTTFLFLVAGVPKSERTTVHKNLTFSLASAEGFLMTSEWAKANEACVAVTVAMHFLFL
VAFSWMLVEGLLLWRKVVAVSMHPGPGMRLYHATGWGVPVGIVAVTLAMLPHDYVAPGHC
WLVHTNAIWAFVGPVLEFVLTVS

Figure 5B

GGTGCCTGGGCCACCACAGGCTGCTCCGTGGCTGCCCTGTACCTGGACTCCACCGCCTGC
TTCTGCAACCACAGCACCAGCTTTGCCATCCTGCTGCAAATCTATGAAGTACAGAGAGGC
CCTGAGGAGGAGTCGCTGCTGAGGACTCTGTCAATTTGTGGGCTGTGGCGTGTCTTCTGC
GCCCTCACCACCACCTTCTTGCTCTTCCTGGTGGCCGGGGTCCCCAAGTCAGAGCGAACC
ACAGTCCACAAGAACCTCACCTTCTCCCTGGCCTCTGCCGAGGGCTTCCTCATGACCAGC
GAGTGGGCCAAGGCCAATGAGGTGGCATGTGTGGCTGTACAGTCGCAATGCACTTCCTC
TTTCTGGTGGCATTCTCCTGGATGCTGGTGGAGGGGCTGCTGCTGTGGAGGAAGGTGGTA
GCTGTGAGCATGCACCCGGGCCCAGGCATGCGGCTCTACCACGCCACAGGCTGGGGCGTG
CCTGTGGGCATCGTGGCGGTACCCCTGGCCATGCTCCCCATGACTACGTGGCCCCCGGA
CATTGCTGGCTCAATGTGCACACAAATGCCATCTGGGCCTTCGTGGGGCCTGTGCTCTTC
GTGCTGACTGTGAGC

"000004" ctg12789

Figure 6

Sequence Name:

GW.A.ctg12776-000000.33.0

Figure 6A

MKSPRRTTLCMLFIVIIYSSKAALNWNYESITHPLLHEHEPAGEEALRQKRAVATKSPTAE
EYTVNIEISFENASFLDPIKAYLNSLSFPIHGNNTDQITDILSINVTTVCRPAGNEIWCS
CETGYGWPRERCLHNLICQERDVFLPGHHCSCLKELPPNGPFCLLQEDVTLNMRVRLNVG
FQEDLMNTSSALYRSYKTDLETARKGYGILPGFKGVTVTGFKSGSVVVTYEVKTTPPSLE
LIHKANEQVQSLNQTYKMDYNSFQAVTINESNFFVTPEIIFEGDTVSLVCEKEVLSSNV
SWRYEEQQLEIQNSSRFSIYTALENNMTSVSKLTIHNITPGDAGEYVCKLILDI FEYECK
KKIDVMPQIILANEEMKVMCDNNPVSLNCCSQGNVNWSKVEWKQEGKINIPGTPETDIDS
SCSRYTLKADGTQCPSGSSGTTVIYTCEFI SAYGARGSANIKVTFISVANLTITPDPISV
SEGQNFSIKCIDSVSNYDEVYWNTSAGIKIYQRFYTTTRYLDGAESVLTVKTSTREWNGT
YHCIFRYKNSYSIATKDVIVHPLPLKLNIMVDPLEATVSCSGSHHIKCCIEEDGDYKVTF
HTGSSSLPAAKEVNKKQVCYKHNFNASSVSWCSKTVDVCCHTNAANNSVWSPSMKLNLV
PGENITCQDPVIGVGEQKVIQKLCRFSNVPSSPESPIGGTITYKCVGSQWEEKRND CIS
APINSLLOMAKLIKSPSQDEMLPTYLKDLSISIDKAEHEISSSPGSLGAIINILDLLSTV
PTQVNSEMMTVLSTVNVLGKPV LNTWKVLQQQWNTNQSSQLLHSVERFSQALQSGDSPPL
SFSQTNVQMSSMVIKSSHPETYQORFVFPYFDLWGNVVIDKSYLENLQSDSSIIVTMAFPT
LQAILAQDIQENNAESLVMTTTVSHNTTMPFRISMTFKNNSPSGGETKCVFWNFR LANN
TGGWDSSGCVVEEGDGNVTCICDHLTSFSILMSPDSPDPSSLLGILLDIISYVGVGFSI
LSLAACLVVEAVVWKSVTKNRTSYMRHTCIVNIAASLLVANTWFI VVAIQDNRYILCKT
ACVAATFFIHFYLSVFFWMLTLGLMLFYRLV FILHETSRSTQKAI AFCLGYGCPLAISV
ITLGATQPREVYTRKNVCWLNWEDTKALLAFAIPALIIVVVNITITIVITKILRPSIGD
KPKQEKSSLFQISKIGVLTPLLGLTWGFGLTTFVPGTNLVFHIIIFAILNVFQLFILLF
GCLWDLKQEALLNKFSLSRWSSQHSKTSLSGSSTPVFSMSSPISRRFNNLFGKTGTYNVST
PEATSSSSLENNSSSASSLLN

Figure 6B

ATGAAATCCCCAAGGAGAACCACCTTTGTGCCTCATGTTTATTGTGATTTATTCTTCCAAA
GCTGCACTGAACTGGAATTACGAGTCTACTATTATCCTTTGAGTCTTCATGAACATGAA
CCAGCTGGTGAAGAGGCACTGAGGCAAAAACGAGCCGTTGCCACAAAAAGTCCTACGGCT
GAAGAATACACTGTTAATATTGAGATCAGTTTTGAAAATGCATCCTTCCTGGATCCTATC
AAAGCCTACTTGAACAGCCTCAGTTTTCCAATTCATGGGAATAACACTGACCAAATTACC
GACATTTTGAGCATAAATGTGACAACAGTCTGCAGACCTGCTGGAAATGAAATCTGGTGC
TCCTGCGAGACAGGTTATGGGTGGCCTCGGGAAAGGTGTCTTCACAATCTCATTTGTCAA
GAGCGTGACGTCTTCCTCCCAGGGCACCATTGCAGTTGCCTTAAAGAACTGCCTCCCAAT
GGACCTTTTTGCCTGCTTCAGGAAGATGTTACCCTGAACATGAGAGTCAGACTAAATGTA
GGCTTTCAGAAGACCTCATGAACACTTCCTCCGCCCTCTATAGGTCCTACAAGACCGAC
TTGGAACACAGCGTTCCGGAAGGTTACGGAATTTTACCAGGCTTCAAGGGCGTGACTGTG
ACAGGGTTCAAGTCTGGAAGTGTGGTTGTGACATATGAAGTCAAGACTACACCACCATCA
CTTGAGTTAATACATAAAGCCAATGAACAAGTTGTACAGAGCCTCAATCAGACCTACAAA
ATGGACTACAACCTCTTTCAAGCAGTTACTATCAATGAAAGCAATTTCTTTGTACACCA
GAAATCATCTTTGAAGGGGACACAGTCAGTCTGGTGTGTGAAAAGGAAGTTTTGTCTCC
AATGTGTCTTGGCGCTATGAAGAACAGCAGTTGGAATCCAGAACAGCAGCAGATTCTCG
ATTTACACCGCACTTTTCAACAACATGACTTCGGTGTCCAAGCTCACCATCCACAACATC
ACTCCAGGTGATGCAGGTGAATATGTTTGCAAACCTGATATTAGACATTTTTGAATATGAG
TGCAAGAAGAAAATAGATGTTATGCCCATCCAAATTTTGGCAAATGAAGAAATGAAGGTG
ATGTGCGACAACAATCCTGTATCTTTGAACTGCTGCAGTCAGGGTAATGTTAATTGGAGC
AAAGTAGAATGGAAGCAGGAAGGAAAAATAAATATTCCAGGAACCCCTGAGACAGACATA
GATTCTAGCTGCAGCAGATACACCCTCAAGGCTGATGGAACCCAGTGCCCAAGCGGGTCG

TCTGGAACAACAGTCATCTACACTTGTGAGTTCATCAGTGCCTATGGAGCCAGAGGCAGT
GCAAACATAAAAGTGACATTTCATCTCTGTGGCCAATCTAACAAATAACCCCGGACCCAATT
TCTGTTTTCTGAGGGACAAAACCTTTTCTATAAAATGCATCAGTGATGTGAGTAACTATGAT
GAGGTTTTATTGGAACACTTCTGCTGGAATTTAAATATACCAAAGATTTTATACCACGAGG
AGGTATCTTGATGGAGCAGAATCAGTACTGACAGTCAAGACCTCGACCAGGGAGTGGAAT
GGAACCTATCACTGCATATTTAGATATAAGAATTCATACAGTATTGCAACCAAAGACGTC
ATTGTTACCCCGCTGCCCTCTAAAGCTGAACATCATGGTTGATCCTTTGGAAGCTACTGTT
TCATGCAGTGGTTCCTCATCATCAAGTGCTGCATAGAGGAGGATGGAGACTACAAAGTT
ACTTTCCATACGGGTTCCCTCATCCCTTCCTGCTGCAAAAGAAGTTAACAAAAACAAGTG
TGCTACAAACACAATTTCAATGCAAGCTCAGTTTCCTGGTGTTCAAAAACCTGTTGATGTG
TGTGTCACTTTTACCAATGTCTAATAATTCAGTCTGGAGCCCATCTATGAAGCTGAAT
CTGGTTTCCTGGGGAAAAACATCACATGCCAGGATCCCGTAATAGGTGTCTGGAGAGCCGGGG
AAAGTCATCCAGAAGCTATGCCGGTTCCTCAAACGTTCCCGAGCAGCCCTGAGAGTCCCAT
GGCGGGACCATCACTTACAAATGTGTAGGCTCCAGTGAGGAGGAGAAGAGAAATGACTGC
ATCTCTGCCCCAATAAACAGTCTGCTCCAGATGGCTAAGGCTTTGATCAAGAGCCCCTCT
CAGGATGAGATGCTCCCTACATACCTGAAGGATCTTTCTATTAGCATAGACAAAGCGGAA
CATGAAATCAGCTCTTCTCCTGGGAGTCTGGGAGCCATTATTAACATCCTTGATCTGCTC
TCAACAGTTCGAACCCCAAGTAAATTCAGAAATGATGACGCACGTGCTCTCTACGGTTAAT
GTCATCCTTGGAAGCCCGTCTTGAACACCTGGAAGGTTTTACAACAGCAATGGACCAAT
CAGAGTTCACAGCTACTACATTCAGTGGAAAGATTTTCCCAAGCATTACAGTCGGGAGAT
AGCCCTCCTTTGCTCTTCTCCCAAACTAATGTGCAGATGAGCAGCATGGTAATCAAGTCC
AGCCACCCAGAAACCTATCAACAGAGGTTTGTCTTCCCATACTTTGACCTCTGGGGCAAT
GTGGTCATTGACAAGAGCTATCTAGAAAACCTTGCAAGTCCGATTTCGTCTATTGTCACCATG
GCTTTCCCAACTCTCCAAGCCATCCTTGCCAGGATATCCAGGAAAATAACTTTGCAGAG
AGCTTAGTGATGACAACCACTGTCAGCCACAATACAACTATGCCATTACAGGATTTCAATG
ACTTTTAAAGAACATAGCCCTTCAGGCGGCGAAACGAAGTGTGTCTTCTGGAACCTCAGG
CTTGCCAAACACAGGGGGGTGGGACAGCAGTGGGTGCTATGTAGAAGAAGTGATGGG
GACAATGTACCTGTATCTGTGACCACCTAACATCATTCTCCATCCTCATGTCCCCTGAC
TCCCAGATCCTAGTTCTCTCCTGGGAATACTCCTGGATATTATTTCTTATGTTGGGGTG
GGCTTTTCCATCTTGAGCTTGCGCAGCCTGTCTAGTTGTGGAAGCTGTGGTGTGGAATCG
GTGACCAAGAACCAGGACTTCTTATATGCGCCACACCTGCATAGTGAATATCGCTGCCTCC
CTTCTGGTCGCCAACACCTGGTTTCATTGTGGTCTGCTGCCATCCAGGACAATCGCTACATA
CTCTGCAAGACAGCCTGTGTGGCTGCCACCTTCTTCATCCACTTCTTCTACCTCAGCGTC
TTCTTCTGGATGCTGACACTGGGCCTCATGCTGTTCTATCGCCTGGTTTTTCATTCTGCAT
GAAACAAGCAGGTCCACTCAGAAAGCCATTGCCTTCTGTCTTGGCTATGGCTGCCCCTT
GCCATCTCGGTCATCACGCTGGGAGCCACCCAGCCCCGGGAAGTCTATACGAGGAAGAAT
GTCTGTTGGCTCAACTGGGAGGACACCAAGGCCCTGCTGGCTTTTCGCCATCCCAGCACTG
ATCATTGTGGTGGTGAACATAACCATCACTATTGTGGTCATACCAAGATCCTGAGGCCT
TCCATTGGAGACAAGCCATGCAAGCAGGAGAAGAGCAGCCTGTTTCAGATCAGCAAGAGC
ATTGGGGTCCCTCACACCACTCTTGGGCCTCACTTGGGGTTTTGGTCTCACCCTGTGTTT
CCAGGGACCAACCTTGTGTTCCATATCATATTTGCCATCCTCAATGTCTTCCAGGGATTA
TTCATTTTACTCTTTGGATGCCTCTGGGATCTGAAGGTACAGGAAGCTTTGCTGAATAAG
TTTTTATTGTCGAGATGGTCTTCACAGCACTCAAAGTCAACATCCCTGGGTTCATCCACA
CCTGTGTTTTCTATGAGTCTCCAATATCAAGGAGATTTAACAAATTTGTTTGGTAAAACA
GGAACGTATAATGTTTCCACCCCAAGCAACCAGCTCATCCCTGGAAAACCTCATCCAGT
GCTTCTTCGTTGCTCAAC

Figure 8

Sequence Name:

GW.S.ctg16790-000000.13.0

Figure 8A

GTTGDWSSEGCSTEVRPEGTVCCCDHLTFFALLLPDQSTVHILTRISQAGCGVSMIFL
AFTIILYAFLRLSRERFKSEDAPKIHVALGGSFLFLNLAFLVNVGSGSKGSDAACWARGA
VFHYFLLCAFTWMGLEAFHLYLLAVRVFNTYFGHYFLKLSLVGWGLPALMVI GTGSANSY
GLYTIRDRENRTSLELCWFREGTTMYALYITVHGYFLITFLFGMVVLALVVKIFTL SRA
TAVKERGKNRKVLTLGLSSLVGV TWGLAIFTPLGLSTVYIFALFNSLQVDFYILIFY

Figure 8B

GGGACCACTGGAGACTGGTCTTCTGAGGGCTGCTCCACGGAGGTCAGACCTGAGGGGACC
GTGTGCTGCTGTGACCACCTGACCTTTTTCGCCCTGCTCCTGAGACCCACCTTGGACCAG
TCCACGGTG CATATCCTCACACGCATCTCCCAGGCGGGCTGTGGGGTCTCCATGATCTTC
CTGGCCTTCACCATATTCTTTATGCCTTCTGAGGCTTCCCGGGAGAGGTTCAAGTCA
GAAGATGCCCCAAAGATCCACGTGGCCCTGGGTGGCAGCCTGTTCCCTCCTGAATCTGGCC
TTCTTGGTCAATGTGGGGAGTGGCTCAAAGGGGTCTGATGCTGCCTGCTGGGCCCCGGGG
GCTGTCTTCCACTACTTCTGCTCTGTGCCTTCACCTGGATGGGCCTTGAAGCCTTCCAC
CTCTACCTGCTCGCTGT CAGGGTCTTCAACACCTACTTCGGGCACTACTTCTGAAGCTG
AGCCTGGTGGGCTGGGGCCTGCCCCCCTGATGGTCATCGGCACTGGGAGTGCCAACAGC
TACGGCCTCTACACCATCCGTGATAGGGAGAACCGCACCTCTCTGGAGCTATGCTGGTTC
CGTGAAGGGACAACCATGTACGCCCTCTATATCACCGTCCACGGCTACTTCTCATCACC
TTCCTCTTTGGCATGGTGGTCTTGGCCCTGGTGGTCTGGAAGATCTTACCCTGTCCCGT
GCTACAGCGGTCAAGGAGCGGGGGAAGAACC GGAAGAAGGTGCTCACCCTGCTGGGCCTC
TCGAGCCTGGTGGGTGTGACATGGGGGTGGCCATCTTCACCCCGTTGGGCCTCTCCACC
GTCTACATCTTTGCACTTTTCAACTCCTTGCAAGTTGATTTTTACATATTGATCTTCTAT

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Figure 9

Sequence Name:

GW.S.ctg12776-000000.172.0

Figure 9A

NHILDTAAISNWFIPNKNASSDLLQSVNLFARQLHIHNNSENIVNELFIQTKGFHINHN
TSEKSLNFSMSMNNTTEDILGMVQIPRQELRKLWPNASQAISIAFPTLGAILREAHLQNV
SLPRQVNLVLSVVLPERLQEIILTFEKINKTRNARAQCVGWHSKKRWDEKACQMMLDI
RNEVKKRCNYTSVVMFSILMSSKSMTDKVLDYITCIGLSVSILSLVLCIIIEATVWSRV
VVTEISYMRHVCIVNIAVSLLTANVWFIIGSHFNKAQDYNMCVAVTFFSHFFYLSLFFW
MLFKALLIIYGILVIFRRMMKSRMMVIGFAIGYGCPLIIAVTTVAITEPEKGYIRPEACW
LNWDNTKALXAFaipafvIVAVNLIVLVAVNTQRPSIGSSKSQDVVIIMRISKNVAIL
TPLLLGLTWGFGIATLIEGTSLTFFHIIIFALLNAFQFFILLFGTIMDHKV

Figure 9B

AACCACATCCTCGACACAGCAGCCATTTCAAACCTGGGCTTTCATTCCCAACAAAAATGCC
AGCTCGGATTTGTTGCAGTCAGTGAATTTGTTGCCAGACAACCTCCACATCCACAATAAT
TCTGAGAACATTGTGAATGAACCTTTCATTGACACAAAAGGGTTTCACATCAACCATAAT
ACCTCAGAGAAAAGCCTCAATTTCTCCATGAGCATGAACAATACCACAGAAGATATCTTA
GGAATGGTACAGATTCCCAGGCAAGAGCTAAGGAAGCTGTGGCCAAATGCATCCCAAGCC
ATTAGCATAGCTTTCCCAACCTTGGGGGCTATCCTGAGAGAAGCCCACTTGCAAAATGTG
AGTCTTCCCAGACAGGTAAATGGTCTGGTGCTATCAGTGGTTTTACCAGAAAGGTTGCAA
GAAATCATACTCACCTTCGAAAAGATCAATAAAAACCCGCAATGCCAGAGCCCAGTGTGTT
GGCTGGCACTCCAAGAAAAGGAGATGGGATGAGAAAAGCGTGCCAAATGATGTTGGATATC
AGGAACGAAGTGAAATGCCGCTGTAACCTACACCAGTGTGGTGATGTCTTTTTCCATTCTC
ATGTCTCCAAATCGATGACCGACAAAGTTCTGGACTACATCACCTGCATTGGGCTCAGC
GTCTCAATCCTAAGCTTGGTTCTTGCCTGATCATTGAAGCCACAGTGTGGTCCCGGGTG
GTTGTGACGGAGATATCATACATGCGTCACGTGTGCATCGTGAATATAGCAGTGTCCCTT
CTGACTGCCAATGTGTGGTTTATCATAGGCTCTCACTTTAACATTAAGGCCCAGGACTAC
AACATGTGTGTTGCAGTGACATTTTTCAGCCACTTTTCTACCTCTCTCTGTTTTTCTGG
ATGCTCTTCAAAGCATTGCTCATCATTTATGGAATATTGGTCATTTTCCGTAGGATGATG
AAGTCCCGAATGATGGTCATTGGCTTTGCCATTGGCTATGGGTGCCCATTGATCATTGCT
GTCACTACAGTTGCTATCACAGAGCCAGAGAAAAGGCTACATAAGACCTGAGGCCTGTTGG
CTTAACCTGGGACAATACCAAAGCCCTTTAAGCATTGTCATCCCGGCGTTCGTGATTGTG
GCTGTAAATCTGATTGTGGTTTTGGTTGTTGCTGTCAACACTCAGAGGCCCTCTATTGGC
AGTTCCAAGTCTCAGGATGTGGTCATAATTATGAGGATCAGCAAAAATGTTGCCATCCTC
ACTCCACTGCTGGGACTGACCTGGGGTTTTGGAATAGCCACTCTCATAGAAGGCACTTCC
TTGACGTTCCATATAATTTTTGCCTTGCTCAATGCTTTCCAGGGTTTTTTCATCCTGCTG
TTTGAACCATATGGATCACAAGGTA

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